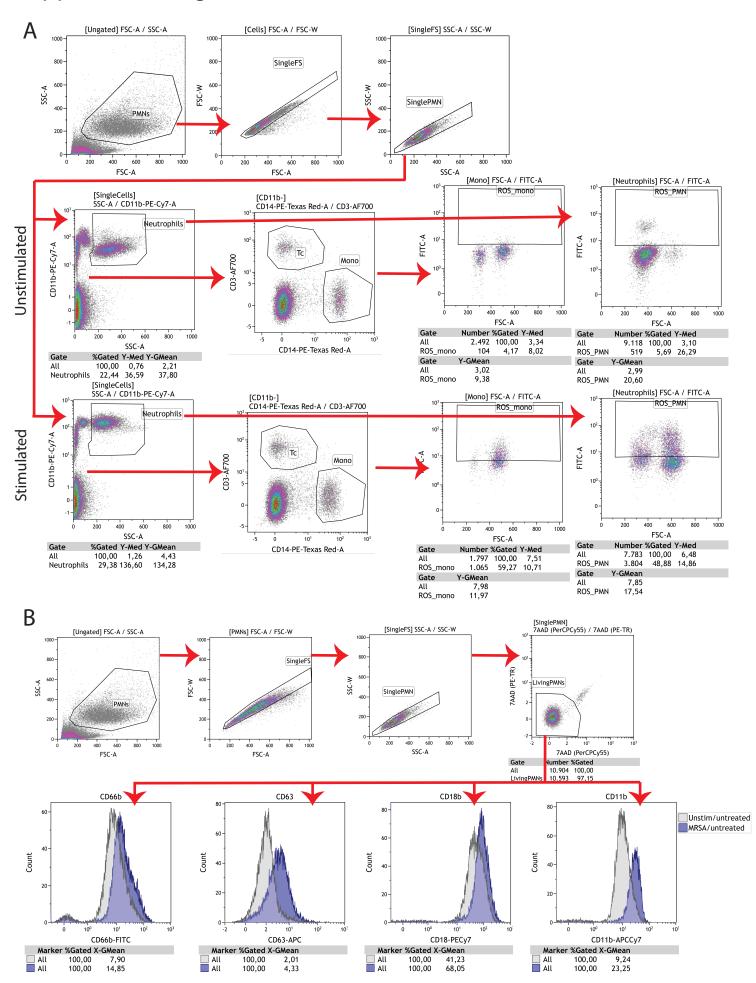
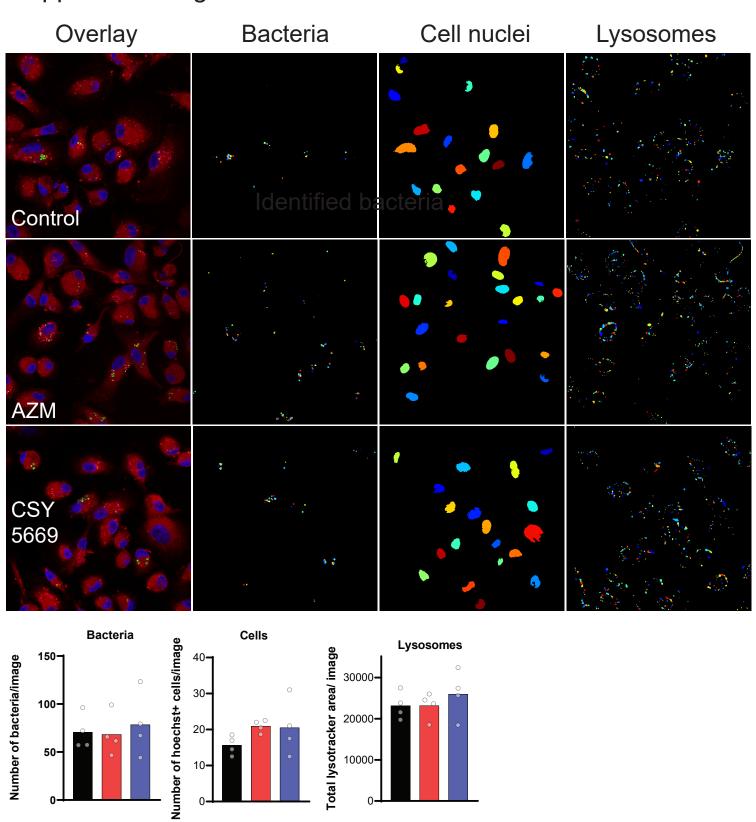


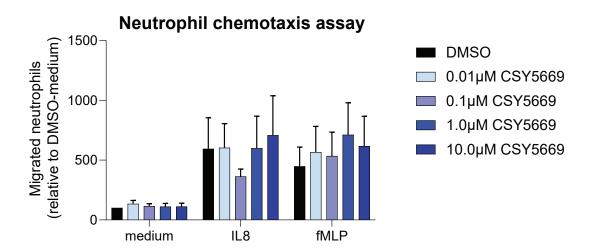
Supplemental figure 1: gating strategy as used to determine cell influx into BALF of MRSA-infected mice. Beads were used to quantify absolute cell influx and were fluorescent in all channels.



Supplemental figure 2: gating strategies as used in a full blood ROS assay (A) and neutrophil degranulation assay (B). In both unstimulated and MRSA-stimulated samples were included.



Supplemental figure 3: quantification of bacterial colocalization with lysosomes. Macrophages were infected with GFP-expressing MRSA (green), treated with CSY5669, AZM or control. After 120 min cells cells were stained withy lysotracker (red) and Hoechst nuclear stain (blue) (A). Each condition was performed in triplicate and three images were obtained per sample (i.e. 9 pictures / condition) and presented with one dot in quantification plots. Representative image displaying automated quantification using cellprofiler of bacteria, (cellular) nuclei and lysotracker+ organelles, wherein every individual identified object is presented with a unique color (A). Quantification of number of bacteria, number of cells and total area of lysotracker events is presented. The level of bacterial colocalization and number of lysotracker events identified per cell is presented in Figure 4H.



Supplemental figure 4: CSY5669 does not affect neutrophil chemotaxis in vitro. Isolated neutrophils were treated with 0.01-10uM CSY5669 or DMSO and seeded in a transwell insert to determine chemotaxis towards medium, IL-8 or N-formylmethionyl-leucyl-phenylalanine (fMLP) after 1 hour (n=3). Data are presented as mean ± SD. Statistical relevance was tested using one-way ANOVA.